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ATTY DOCKET NO: P-SH 4810

SERIAL NO: 09/882,355

APPLICANT: Shankar Subramaniam

INFORMATIONAL DISCLOSURE STATEMENT BY APPLICANT

FILING DATE: June 14, 2001

GROUP: 1645

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U. S. PATENT DOCUMENTS

EXAM. INITIALS	DOCUMENT NUMBER	DATE	NAME	CLASS	SUB-CLASS	FILING DATE
gw	5,143,854	09/01/92	Pirrung et al.			
	5,489,678	02/06/96	Fodor et al.			
	5,571,639	11/05/96	Hubbell et al.			
	5,667,195	09/16/97	McCormick			
	5,744,101	04/28/98	Fodor et al.			

FOREIGN PATENT DOCUMENTS

EXAM. INITIALS	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB-CLASS	TRANSLATION (YES/NO)
gw	EP1026258A2	08/09/00	Europe			

OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages)

gw	Adams et al., "The Genome Sequence of <i>Drosophila Melanogaster</i> ," <u>Science</u> , 287:2185-2195 (2000).
gw	Arsilan et al., "A New Approach to Sequence Comparison: Normalized Sequence Alignment," <u>Bioinformatics</u> , 17(4):327-337 (2001).

EXAMINER <i>gw</i>	DATE CONSIDERED 9/28/02
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<i>fw</i>	C. Elegans Sequencing Consortium, "Genome Sequence of the Nematode C. Elegans: A Platform for Investigating Biology," <u>Science</u> , 282:2012-2018 (1998).
	Cheung et al., "Linkage-Disequilibrium Mapping Without Genotyping," <u>Nat. Gen.</u> , 18:225-230 (1998).
	Hayward et al., "Shotgun DNA Microarrays and Stage-Specific Gene Expression in Plasmodium Falciparum Malaria," <u>Mol. Microbiol.</u> , 35(1):6-14 (2000).
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	Mironov et al., "Performance-Guarantee Gene Predictions Via Spliced Alignment," <u>Genomics</u> , 51:332-339 (1998).
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	Selinger et al., "RNA Expression Analysis Using a 30 Base Pair Resolution Escherichia Coli Genome Array," <u>Nat. Biotech.</u> , 18:1262-1268 (2000).
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	Sze et al., "Algorithms and Software for Support of Gene Identification Experiments," <u>Bioinformatics</u> , 14(1):14-19 (1998).
	Venter et al., "Shotgun Sequencing of the Human Genome," <u>Science</u> , 280:1540-1542 (1998).
<i>fw</i>	Venter et al., "The Sequence of the Human Genome," <u>Science</u> , 291:1304-1351 (2001).

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